

## **Product datasheet for SC200655**

## DPM3 (NM 153741) Human 3' UTR Clone

## **Product data:**

**Product Type:** 3' UTR Clones

Product Name: DPM3 (NM\_153741) Human 3' UTR Clone

**Vector:** pMirTarget (PS100062)

Symbol: DPM3

Synonyms: CDG10; MDDGB15; MDDGC15

**ACCN:** NM\_153741

**Insert Size:** 101 bp

Insert Sequence: >SC200655 3'UTR clone of NM\_153741

The sequence shown below is from the reference sequence of NM\_153741. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

GACTTAGCCCGCAGGGGGCTGCGCTTCTGACAGCCTAACCCCATTCCTGTGCGGACAGCCCTTCCTCCC

ATTTCCCATTAAAGAGCCAGTTTATTTTCTAA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

**OTI Disclaimer:** Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the

reference sequence but may contain minor differences, e.g., single nucleotide

polymorphisms (SNPs).

**Components:** The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

**RefSeg:** NM 153741.2



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## DPM3 (NM\_153741) Human 3' UTR Clone - SC200655

Summary: Dolichol-phosphate mannose (Dol-P-Man) serves as a donor of mannosyl residues on the

lumenal side of the endoplasmic reticulum (ER). Lack of Dol-P-Man results in defective surface expression of GPI-anchored proteins. Dol-P-Man is synthesized from GDP-mannose and dolichol-phosphate on the cytosolic side of the ER by the enzyme dolichyl-phosphate mannosyltransferase. The protein encoded by this gene is a subunit of dolichyl-phosphate

mannosyltransferase and acts as a stabilizer subunit of the dolichyl-phosphate

mannosyltransferase complex. [provided by RefSeq, Jul 2008]

**Locus ID:** 54344

MW: 3.7